Local DAG for enriched terms in GO Molecular Function

Nodes sized according to Binomial Fold Enrichment

SH3 domain binding

2.18

tumor necrosis factor receptor superfamily binding

2.59

tumor necrosis factor receptor binding

2.85

phosphoric diester hydrolase activity

2.38

non-membrane spanning protein tyrosine kinase activity

Node size legend: < 1   2   3   4   > 5